

PhosFate analysis

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 An abbreviated version of this protocol was published in eLIFE in Jan 2020

Dynamic post-translational modification profiling of *Mycobacterium tuberculosis*-infected primary macrophages

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Detailed protocol

The input list for Phosfate was generated by mapping all of the phosphopeptide changes from the mouse to the human proteome using a R script.

How to cite: (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Budzik, J. M. and Cox, J. S.(2021). PhosFate analysis. Bio-protocol Preprint. bio-protocol.org/prep1063.
2. Budzik, J. M., Swaney, D. L., Jimenez-Morales, D., Johnson, J. R., Garelis, N. E., Repasy, T., Roberts, A. W., Popov, L. M., Parry, T. J., Pratt, D., Ideker, T., Krogan, N. J. and Cox, J. S.(2020). Dynamic post-translational modification profiling of *Mycobacterium tuberculosis*-infected primary macrophages. eLIFE. DOI: [10.7554/eLife.51461](https://doi.org/10.7554/eLife.51461)

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